

<u>DB Name</u>	<u>Query</u>	<u>Hit Count</u>	<u>Set Name</u>
USPT	114 same brain	0	<u>L23</u>
USPT	19 same brain	33	<u>L22</u>
USPT	19 same disease\$	7	<u>L21</u>
USPT	110 same disease\$	0	<u>L20</u>
USPT	114 same disease\$	0	<u>L19</u>
USPT	115 same disease\$	0	<u>L18</u>
USPT	115 same (human or non)	4	<u>L17</u>
USPT	114 same (human or non)	4	<u>L16</u>
USPT	114 same (tissue\$ or cell\$)	9	<u>L15</u>
USPT	110 same (double near0 strand\$)	12	<u>L14</u>
USPT	18 same (driver near0 normaliz\$)	0	<u>L13</u>
USPT	19 same (driver near0 normaliz\$)	0	<u>L12</u>
USPT	110 same (driver near0 normaliz\$)	0	<u>L11</u>
USPT	19 same mRNA	138	<u>L10</u>
USPT	18 same (cDNA near0 librar\$)	460	<u>L9</u>
USPT	select\$ near0 clone\$	4881	<u>L8</u>
USPT	mak\$ same (subtract\$ near0 cDNA near0 library)	3	<u>L7</u>
USPT	compar\$ same (subtract\$ near0 cDNA near0 library)	19	<u>L6</u>
USPT	transcriptome	0	<u>L5</u>
USPT	13 same (advantag\$ or useful\$)	2	<u>L4</u>
USPT	identif\$ same (redundant near0 clone\$)	8	<u>L3</u>
USPT	11 same (redundant near0 clone\$)	1	<u>L2</u>
USPT	cDNA near0 librar\$	9330	<u>L1</u>